

Repeated Observations

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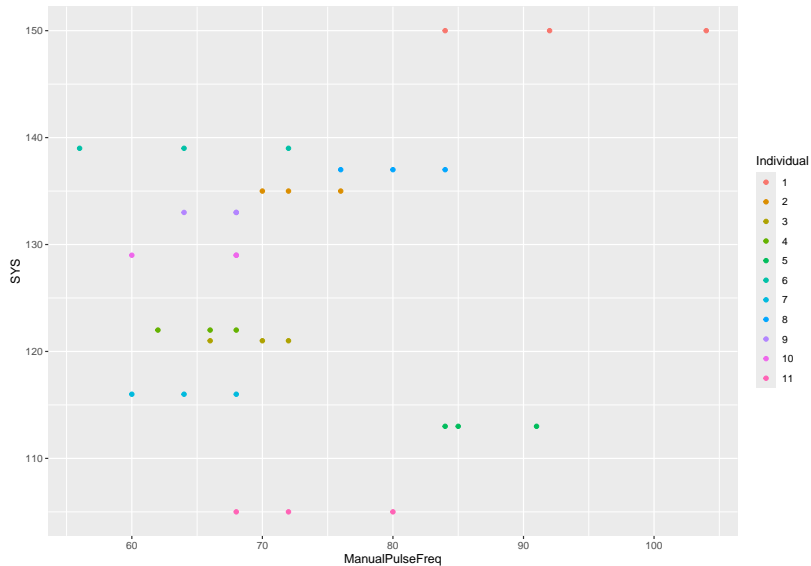
2024-05-06

Three Types of Datastructures

1. Repeated observations in the predictors
2. Repeated observations in the response variables
3. Repeated observations in both the predictors and the responses

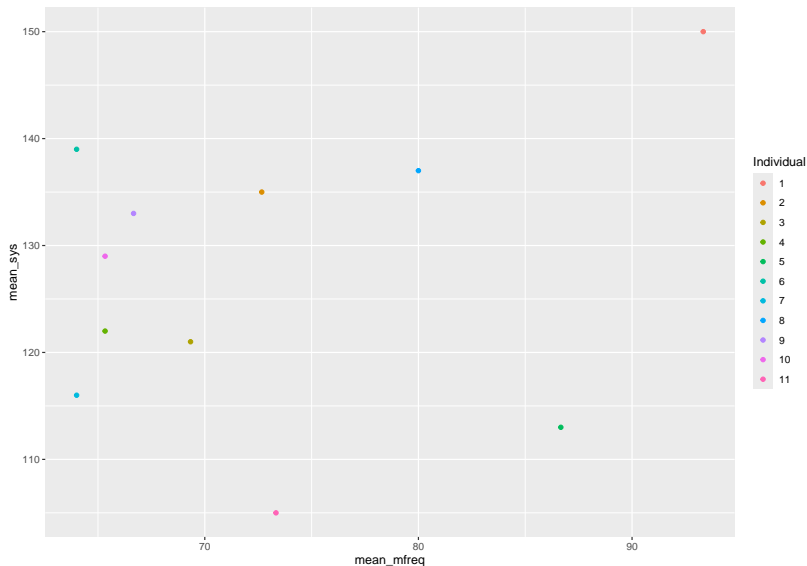
Repeated Observations in the Predictors

Dataset on blood pressure and heart frequency



Statistical Analysis

- ▶ Linear regression of response on mean (median) of predictors
- ▶ Example: SYS as response



Regression

```
lm_msys_mfreq <- lm(mean_sys ~ mean_mfreq, data = tbl_bp_mfreq)
summary(lm_msys_mfreq)
```

Call:

```
lm(formula = mean_sys ~ mean_mfreq, data = tbl_bp_mfreq_mea
```

Residuals:

Min	1Q	Median	3Q	Max
-22.459	-6.685	4.270	7.792	15.720

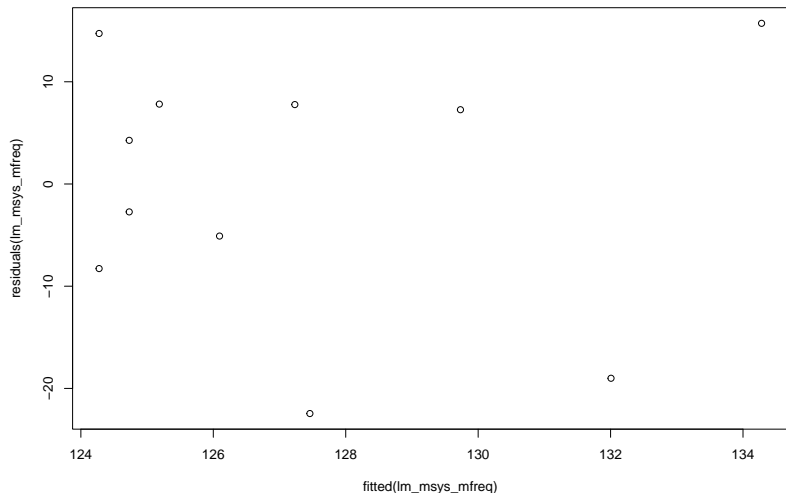
Coefficients:

	Estimate	Std. Error	t value	Pr(> t)	
(Intercept)	102.4456	31.4616	3.256	0.0099	**
mean_mfreq	0.3411	0.4286	0.796	0.4467	

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

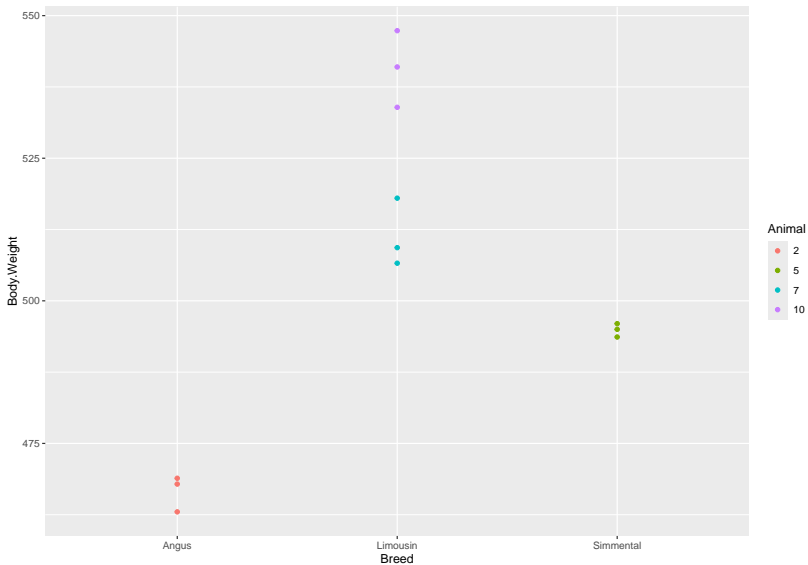
Residuals Plot

```
plot(fitted(lm_msys_mfreq), residuals(lm_msys_mfreq))
```



Repeated observations in the Response

Breed on Body Weight in beef cattle animals



Statistical Analysis

- ▶ Mixed effects model
- ▶ Use `lme4::lmer()` for analysis

```
library(lme4)
lmer_bw_br <- lmer(Body.Weight ~ Breed + (1 | Animal),
                  data = df_bw_br_rep)
```


Results

```
summary(lmer_bw_br)
```

```
Linear mixed model fit by REML ['lmerMod']  
Formula: Body.Weight ~ Breed + (1 | Animal)  
Data: df_bw_br_rep
```

```
REML criterion at convergence: 61.8
```

```
Scaled residuals:
```

Min	1Q	Median	3Q	Max
-1.3714	-0.5383	0.0640	0.3213	1.4305

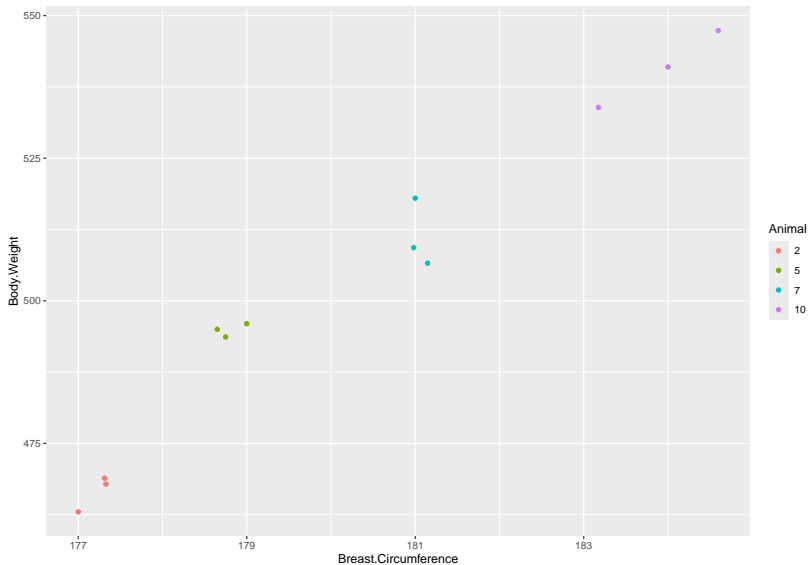
```
Random effects:
```

Groups	Name	Variance	Std.Dev.
Animal	(Intercept)	426.21	20.645
Residual		22.98	4.794

```
Number of obs: 12, groups: Animal, 4
```

Repeated Observations in Both Predictors and Response

Repeated observations in Body Weight and Breast Circumference



Statistical Analysis

- ▶ Use Machine Learning Approaches
- ▶ First approximation:
 - ▶ Compute mean/median of BC
 - ▶ Fit repeated observation mixed model as in type 2